

Supplementary Material: Random Projection for fast and efficient multivariate correlation analysis of high-dimensional data: A new approach

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1 SUPPLEMENTARY TABLES

Table 1. Average PLSC and PLSC-RP weights for high-dimensional neuroimaging data
The table shows average PLSC weights and average PLSC-RP weights for causal voxels and causal SNPs as compared to non-causal voxels and non-causal SNPs. Causal voxels and causal SNPs receive higher weights than non-causal voxels and SNPs. Average weights are very similar for PLSC and PLSC-RP.

| dimensionality of MRI data | PLS analysis | $ \bar{w}_{\text{MRI}} $ for causal voxels | $ \bar{w}_{\text{MRI}} $ for non-causal voxels | $ \bar{w}_{\text{SNP}} $ for causal SNPs | $ \bar{w}_{\text{SNP}} $ for non-causal SNPs |
|-------------------------------|--------------|---|---|---|---|
| 1,000 | PLSC | 0.0503 | 0.0256 | 0.3395 | 0.0976 |
| | PLSC-RP | 0.0506 | 0.0255 | 0.3314 | 0.0995 |
| 10,000 | PLSC | 0.0152 | 0.0078 | 0.3863 | 0.0901 |
| | PLSC-RP | 0.0152 | 0.0078 | 0.3710 | 0.0911 |
| 20,000 | PLSC | 0.0116 | 0.0054 | 0.3440 | 0.0934 |
| | PLSC-RP | 0.0114 | 0.0055 | 0.3266 | 0.0919 |
| 30,000 | PLSC | 0.0104 | 0.0044 | 0.3877 | 0.0881 |
| | PLSC-RP | 0.0104 | 0.0045 | 0.3856 | 0.0898 |
| 40,000 | PLSC | 0.0089 | 0.0041 | 0.3663 | 0.0928 |
| | PLSC-RP | 0.0086 | 0.0041 | 0.3533 | 0.0937 |
| 50,000 | PLSC | 0.0084 | 0.0039 | 0.3143 | 0.0993 |
| | PLSC-RP | 0.0086 | 0.0039 | 0.3123 | 0.1019 |
| 70,000 | PLSC | 0.0061 | 0.0033 | 0.3325 | 0.1005 |
| | PLSC-RP | 0.0058 | 0.0034 | 0.3226 | 0.0973 |
| 90,000 | PLSC | 0.0058 | 0.0029 | 0.2873 | 0.1074 |
| | PLSC-RP | 0.0057 | 0.0029 | 0.2772 | 0.1091 |

Table 2. Average PLSC and PLSC-RP weights for the fMRI face-matching task

Average weights for causal and non-causal voxels and SNPs are very similar for PLSC and PLSC-RP.

| PLS analysis | $ \bar{w}_{\text{MRI}} $ for causal voxels | $ \bar{w}_{\text{MRI}} $ for non-causal voxels | $ \bar{w}_{\text{SNP}} $ for causal SNPs | $ \bar{w}_{\text{SNP}} $ for non-causal SNPs |
|--------------|--|--|--|--|
| PLSC | 0.0059 | 0.0017 | 0.5768 | 0.0296 |
| PLSC-RP | 0.0059 | 0.0017 | 0.5757 | 0.0467 |

Table 3. Average SNP weights for PLSC and PLSC-RP in the Sorbs

The table shows average PLSC weights and average PLSC-RP weights for causal and non-causal SNPs. In addition, it is illustrated how serum vaspin and body height are weighted in the first component of the phenotype weight profile.

| PLS analysis | $ w_{\text{Vaspin}} $ | $ w_{\text{Height}} $ | $ \bar{w}_{\text{SNP}} $ for causal SNPs | $ \bar{w}_{\text{SNP}} $ for non-causal SNPs |
|--------------|-----------------------|-----------------------|--|--|
| PLSC | 0.7068 | 0.0285 | 0.0093 | 0.0013 |
| PLSC-RP | 0.7068 | 0.0294 | 0.0093 | 0.0013 |

Table 4. Average PLSC and PLSC-RP weights for high-dimensional neuroimaging and high-dimensional SNP data

Causal voxels and causal SNPs receive higher weights than non-causal voxels and SNPs. Average weights are very similar for PLSC and PLSC-RP.

| dim. of MRI data | dim. of SNP data | PLS analysis | $ \bar{w}_{\text{MRI}} $ for causal voxels | $ \bar{w}_{\text{MRI}} $ for non-causal voxels | $ \bar{w}_{\text{SNP}} $ for causal SNPs | $ \bar{w}_{\text{SNP}} $ for non-causal SNPs |
|------------------|------------------|--------------|--|--|--|--|
| 1,000 | 1,000 | PLSC | 0.0405 | 0.0288 | 0.0947 | 0.0246 |
| | | PLSC-RP | 0.0435 | 0.0279 | 0.0985 | 0.0245 |
| 10,000 | 10,000 | PLSC | 0.0158 | 0.0077 | 0.0370 | 0.0080 |
| | | PLSC-RP | 0.0151 | 0.0078 | 0.0368 | 0.0079 |
| 20,000 | 20,000 | PLSC | 0.0116 | 0.0054 | 0.0256 | 0.0056 |
| | | PLSC-RP | 0.0106 | 0.0056 | 0.0240 | 0.0056 |
| 40,000 | 40,000 | PLSC | 0.0092 | 0.0041 | 0.0179 | 0.0040 |
| | | PLSC-RP | 0.0092 | 0.0040 | 0.0187 | 0.0040 |
| 50,000 | 1,000 | PLSC | 0.0078 | 0.0040 | 0.1073 | 0.0244 |
| | | PLSC-RP | 0.0070 | 0.0039 | 0.0952 | 0.0246 |
| 1,000 | 50,000 | PLSC | 0.0395 | 0.0290 | 0.0150 | 0.0036 |
| | | PLSC-RP | 0.0361 | 0.0293 | 0.0142 | 0.0036 |

2 SUPPLEMENTARY EQUATIONS

PLSC-RP for dimensionality reduction in X_1 OR X_2

For traditional PLSC, SVD is used to decompose the cross-product matrix A of X_1 and X_2 , which are both standardized column-wise, into three matrices:

$$\text{cov}(X_1, X_2) = A = X_1' X_2 = W_1 S W_2'. \quad (1)$$

Assumed that X_1 is high-dimensional, RP transforms X_1 to a lower dimensional space via the following transformation:

$$X_{1\text{RP}} = X_1 \cdot R, \quad (2)$$

where R is a random matrix and $X_{1\text{RP}}$ is the low-dimensional subspace of X_1 with desired lower dimension k . If we perform PLSC to decompose the cross-product matrix of $X_{1\text{RP}}$ and X_2 , we obtain the weights W_2 for data set X_2 , but weights $W_{1\text{RP}}$ for the reduced data set $X_{1\text{RP}}$. To transform the weights $W_{1\text{RP}}$ back to the original space, that is W_1 , we rearrange the equation for the SVD as follows:

Starting point for the rearrangement: the PLSC equation

$$\text{cov}(X_1, X_2) = A = X_1' X_2 = W_1 S W_2'.$$

If we extend both sides of the equation by w_{2i} , we obtain

$$A \cdot w_{2i} = W_1 S W_2' \cdot w_{2i}.$$

Since W_2 is column-wise orthogonal, we have

$$A \cdot w_{2i} = w_{1i} s_i w_{2i}' \cdot w_{2i}.$$

Rearranging yields

$$\frac{1}{s_i} \cdot A \cdot w_{2i} = w_{1i} \cdot w_{2i}' \cdot w_{2i}.$$

Since the L2-norm for a vector a is given by

$$\begin{aligned} |a| &= \sqrt{a_1^2 + a_2^2 + \dots + a_n^2}, \\ |a|^2 &= a_1^2 + a_2^2 + \dots + a_n^2, \end{aligned}$$

we obtain the weights w_{1i} , $i = 1, \dots, p$, $p = \min(k, d_2)$, as follows:

$$w_{1i} = \frac{1}{s_i \cdot |w_{2i}|^2} \cdot A \cdot w_{2i}. \quad (3)$$

PLSC-RP for dimensionality reduction in X_1 AND X_2

Assumed that both X_1 and X_2 are high-dimensional, RP transforms X_1 and X_2 to lower dimensional spaces via the following transformation:

$$\begin{aligned} X_{1RP} &= X_1 \cdot R_1, \\ X_{2RP} &= X_2 \cdot R_2. \end{aligned} \quad (4)$$

If we perform PLSC to decompose the cross-product matrix of X_{1RP} and X_{2RP} , we obtain weights W_{1RP} and W_{2RP} for the low dimensional subspaces. To transform the weights W_{1RP} back to the original space W_1 , we rearrange the equation for the SVD as follows:

Starting point for the rearrangement: the PLSC equation

$$\text{cov}(X_1, X_{2RP}) = W_1 S W_{2RP}'.$$

If we extend both sides of the equation by w_{2RP_i} , we obtain

$$\text{cov}(X_1, X_{2RP}) \cdot w_{2RP_i} = W_1 S W_{2RP}' \cdot w_{2RP_i}.$$

Since W_{2RP} is column-wise orthogonal, we have

$$\text{cov}(X_1, X_{2RP}) \cdot w_{2RP_i} = w_{1_i} s_i w_{2RP_i}' \cdot w_{2RP_i}.$$

Rearranging yields

$$\frac{1}{s_i} \cdot \text{cov}(X_1, X_{2RP}) \cdot w_{2RP_i} = w_{1_i} \cdot w_{2RP_i}' \cdot w_{2RP_i}.$$

Thus, for the weights w_{1_i} , $i = 1, \dots, p$, $p = \min(k_1, k_2)$, we obtain

$$w_{1_i} = \frac{1}{s_i \cdot |w_{2RP_i}|^2} \cdot \text{cov}(X_1, X_{2RP}) \cdot w_{2RP_i}. \quad (5)$$

Following the same logic, the weights W_{2RP} are transformed back to the original space W_2 by rearranging the equation for the SVD as follows:

Starting point for the rearrangement: the PLSC equation

$$\text{cov}(X_{1RP}, X_2) = W_{1RP} S W_2'.$$

If we extend both sides of the equation by w_{1RP_i}' , we obtain

$$w_{1RP_i}' \cdot \text{cov}(X_{1RP}, X_2) = w_{1RP_i}' \cdot W_{1RP} S W_2'.$$

Since $\mathbf{W}_{1\text{RP}}$ is column-wise orthogonal, we have

$$\mathbf{w}'_{1\text{RP}_i} \cdot \text{cov}(\mathbf{X}_{1\text{RP}}, \mathbf{X}_2) = \mathbf{w}'_{1\text{RP}_i} \cdot \mathbf{w}_{1\text{RP}_i} \cdot s_i \cdot \mathbf{w}'_{2_i}.$$

Rearranging yields

$$\frac{1}{s_i} \cdot \mathbf{w}'_{1\text{RP}_i} \cdot \text{cov}(\mathbf{X}_{1\text{RP}}, \mathbf{X}_2) = \mathbf{w}'_{1\text{RP}_i} \cdot \mathbf{w}_{1\text{RP}_i} \cdot \mathbf{w}'_{2_i}.$$

Thus, for the weights \mathbf{w}_{2_i} , $i = 1, \dots, p$, $p = \min(k_1, k_2)$, we obtain

$$\begin{aligned} \mathbf{w}'_{2_i} &= \frac{1}{s_i \cdot |\mathbf{w}_{1\text{RP}_i}|^2} \cdot \mathbf{w}'_{1\text{RP}_i} \cdot \text{cov}(\mathbf{X}_{1\text{RP}}, \mathbf{X}_2), \\ \mathbf{w}_{2_i} &= \frac{1}{s_i \cdot |\mathbf{w}_{1\text{RP}_i}|^2} \cdot (\text{cov}(\mathbf{X}_{1\text{RP}}, \mathbf{X}_2))' \cdot \mathbf{w}_{1\text{RP}_i}. \end{aligned} \tag{6}$$